Genome Search

Genome Search is another way for selecting and browsing genomes in IMG.

Genomes can be searched via implicit attributes using the **Quick Genome Search** box at the top right corner of every IMG page, as shown in Figure 1.

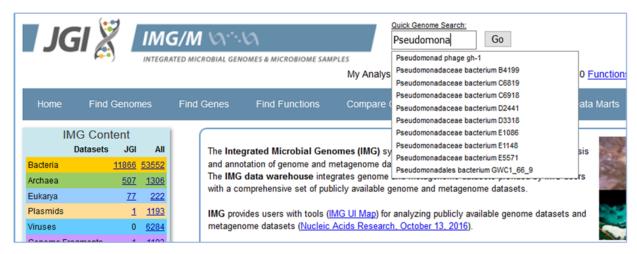


Figure 1. Quick Genome Search.

A detailed search of genomes can be carried out using **Genome Search** on the second-level menu of **Find Genomes**. There are 4 options: by Fields, by Metadata Categories, by Metadata Category Operation, and by Metadata Category Chart. The result of **Genome Search** is a list of genomes, which can be examined individually, or selected and saved for reducing the genome context for future analysis.

By Fields

Genome Search can use *genome fields* as search filter to find genomes, as shown in Figure 2.

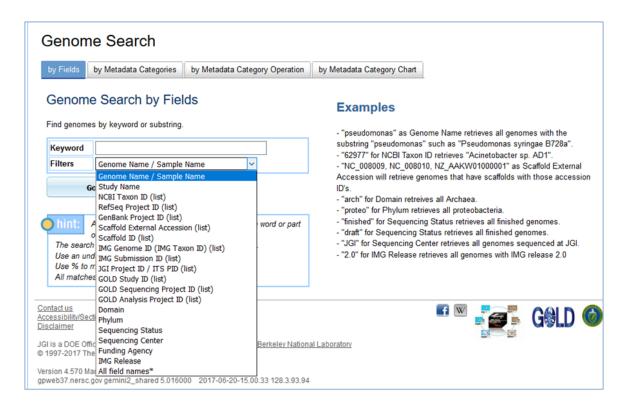


Figure 2. Genome Search by Fields.

Searchable fields are available in the pull-down filter list, including:

- Genome Name / Sample Name
- Study Name
- NCBI Taxon ID (list)
- RefSeq Project ID (list)
- GenBank Project ID (list)
- Scaffold External Accession (list)
- Scaffold ID (list)
- IMG Genome ID (IMG Taxon ID) (list)
- IMG Submission ID (list)
- JGI Project ID / ITS PID (list)
- GOLD Study ID (list)
- GOLD Sequencing Project ID (list)
- GOLD Analysis Project ID (list)
- Domain
- Phylum
- Sequencing Status
- Sequencing Center
- Funding Agency

- IMG Release
- all field names*

Filter options that include "(list)" allow users to enter multiple IDs separated by commas (,). All searches are case insensitive. Check "hint" for more guidance.

By Metadata Categories

Genome search by metadata categories allows users to search genomes using metadata fields from GOLD (see Figure 3). Searchable GOLD metadata categories/fields include:

- Altitude
- Biotic Relationships
- Cell Arrangement
- Cell Shape
- Culture Type
- Uncultured Type
- Depth
- Diseases
- Ecosystem
- Ecosystem Category
- Ecosystem Type
- Ecosystem Subtype
- Energy Source
- Geographic Location
- Gram Staining
- Habitat
- Host Gender
- Isolation
- Metabolism
- Motility
- Oxygen requirement
- Phenotype
- Relevance
- Salinity
- Specific Ecosystem
- Sporulation
- Temperature Range
- Type Strain
- PMO Project Id (list)
- ITS PID (list)
- ITS Proposal Id (list)

GPTS Proposal Id (list)

Filter options that include "(list)" allow users to enter multiple IDs separated by commas (,). All searches are case insensitive. Check "hint" for more guidance.

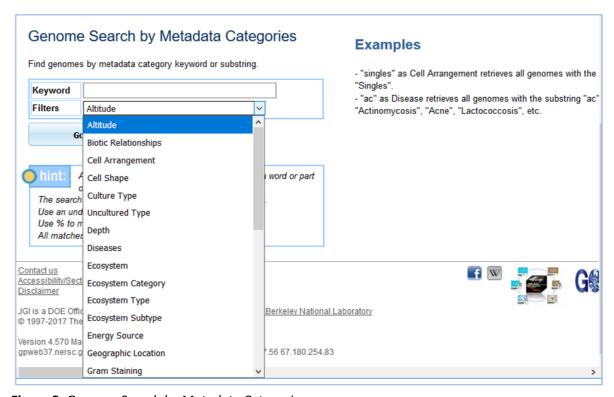


Figure 3. Genome Search by Metadata Categories.

This search option is keyword based. If users do not have idea about what keywords to enter, then they should use the following two options (*metadata category operation* and *metadata category chart*) instead.

By Metadata Category Operation

This option allows users to search multiple conditions on the above mentioned metadata categories using "and" and/or "or" operators. Moreover, users can click on the plus-sign (+) to view pre-defined controlled vocabularies specified for each metadata category. For example, Figure 4 shows that **Biotic Relationships** can be: *Free living, Free-living, endophytic*, or *Symbiotic*. Users can click to select one or more controlled vocabularies. Conditions on multiple values of the same category will be combined using "or" operator.

It is also possible for users to specify conditions on multiple metadata categories (e.g., **Biotic Relationships** and **Cell Shape**). Conditions on different categories will be combined using "and" operator.

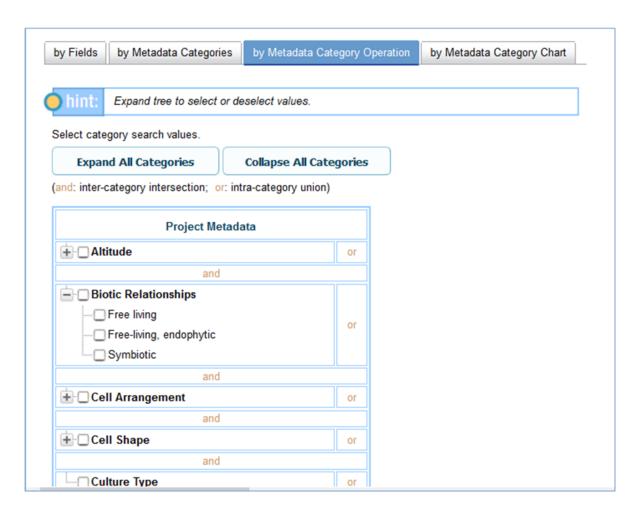


Figure 4. Genome Search by Metadata Category Operation.

It is possible that no genomes will be returned by a metadata category operation search because not all metadata category values have associated genomes in IMG, or there do not exist genomes in IMG satisfying combined search conditions.

To see which GOLD metadata category values actually have associated genomes in IMG, users can use the *metadata category chart* option described immediately below.

By Metadata Category Chart

To search genomes by metadata category chart, click the **by Metadata Category Chart** tab, select a category and then click the **Show Chart** button (see Figure 5(i)). Figure 5 (ii) shows the chart of **Cell Arrangement** and associated genome counts in IMG. Users can click on any bar to view the actual genome list. For example, Figure 5(iii) lists all genomes in IMG with *Tetrads* cell arrangement. Genomes can be selected to add to analysis cart or workspace for further analysis.

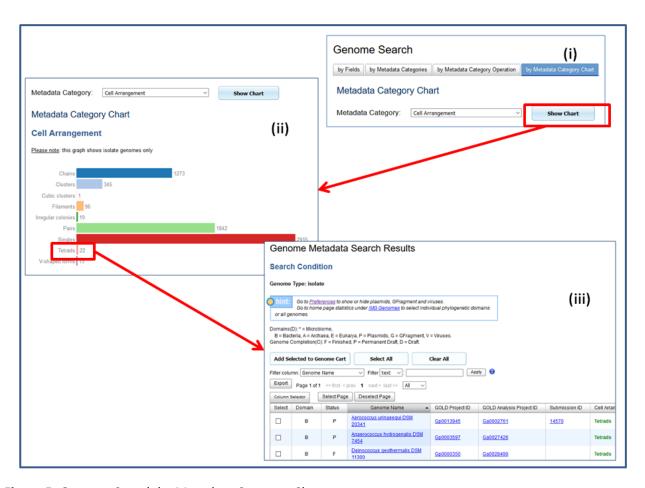


Figure 5. Genome Search by Metadata Category Chart.